

# SEQUENCE LISTING

<110> COGENT NEUROSCIENCE, Inc.  
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<120> COMPOSITIONS AND METHODS FOR DIAGNOSING  
 AND TREATING CONDITIONS, DISORDERS, OR DISEASES INVOLVING  
 CELL DEATH

<130> 10001-005-999

<140> Not Assigned

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<212> DNA

<213> Homo sapiens

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<210> 21  
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<400> 21  
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<211> 93  
<212> DNA  
<213> Homo sapiens

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gaccaccag ccctccggct gctgatgtca tga 93

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<400> 34  
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Thr Ala His Ala Asp His Pro Ala Leu Arg Leu Leu Met Ser  
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<400> 36  
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<400> 37

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gtacagaaga aaaagcgggt ggaccggctg cgccatcacc tgctcccat gtacagctat      180
gacccagctg aggaactgca tgaggctgag caggagctgc tctctgacat gggagacccc    240
aaggtggtac atggctggca gagtggctac cagcacaagc ggatgccact gctggatgtc    300
aagacgtga                                     309

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<210> 38  
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 20          25          30
Val Val Val Ala Val Val Met Tyr Val Gln Lys Lys Lys Arg Val Asp
 35          40          45
Arg Leu Arg His His Leu Leu Pro Met Tyr Ser Tyr Asp Pro Ala Glu
 50          55          60
Glu Leu His Glu Ala Glu Gln Glu Leu Leu Ser Asp Met Gly Asp Pro
 65          70          75          80
Lys Val Val His Gly Trp Gln Ser Gly Tyr Gln His Lys Arg Met Pro
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Leu Leu Asp Val Lys Thr
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<210> 39  
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 <212> DNA  
 <213> Homo sapiens

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tcaccctggt cgggggtggt gtggctgtgg taa                                93

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<210> 40  
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 20          25          30

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<210> 41  
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1				5					10					15	
Leu	Pro	Met	Tyr	Ser	Tyr	Asp	Pro	Ala	Glu	Glu	Leu	His	Glu	Ala	Glu
			20					25					30		
Gln	Glu	Leu	Leu	Ser	Asp	Met	Gly	Asp	Pro	Lys	Val	Val	His	Gly	Trp
		35					40					45			
Gln	Ser	Gly	Tyr	Gln	His	Lys	Arg	Met	Pro	Leu	Leu	Asp	Val	Lys	Thr
	50					55				60					

<211> 480

<213> Homo sapiens

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ccaaggtggt	acatggctgg	cagagtggct	accagcacia	gcggatgcca	ctgctggatg	180
tcaagacgtg	acctgacccc	cttgccccac	ccttcagagc	ctgggggtcct	ggactgcctg	240
gggccctgcc	atctgcttcc	cctgctgtca	cctggctccc	cctgctgggt	gctgggtctc	300
catttctccc	tccaccaccc	ctcagcagca	tctgcttccc	atgccctcac	catcacctca	360
ctgccccag	gccttctgcc	ctttgtgggt	gttgagctca	cgcgccaccc	acaggcactc	420
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<211> 159

<213> Homo sapiens

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1				5					10					15	
Pro	Cys	Thr	Ala	Met	Thr	Gln	Leu	Arg	Asn	Cys	Met	Arg	Leu	Ser	Arg
			20					25					30		
Ser	Cys	Ser	Leu	Thr	Trp	Glu	Thr	Pro	Arg	Trp	Tyr	Met	Ala	Gly	Arg
		35					40					45			
Val	Ala	Thr	Ser	Thr	Ser	Gly	Cys	His	Cys	Trp	Met	Ser	Arg	Arg	Asp
	50					55				60					
Leu	Thr	Pro	Leu	Pro	His	Pro	Ser	Glu	Pro	Gly	Val	Leu	Asp	Cys	Leu
65					70					75					80
Gly	Pro	Cys	His	Leu	Leu	Pro	Leu	Leu	Ser	Pro	Gly	Ser	Pro	Cys	Trp
				85					90					95	
Val	Leu	Gly	Leu	His	Phe	Ser	Leu	His	Pro	Pro	Ser	Ala	Ala	Ser	Ala
			100					105					110		
Ser	His	Ala	Leu	Thr	Ile	Thr	Ser	Leu	Pro	Pro	Gly	Leu	Leu	Pro	Phe
		115					120					125			
Val	Gly	Val	Glu	Leu	Thr	Ala	His	Pro	Gln	Ala	Leu	Ile	Gly	Arg	Gly
	130					135				140					
Phe	Pro	Ser	Gly	Met	Ala	Ala	Ala	Gly	Arg	His	Leu	Cys	Phe	Leu	
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**THE**

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Tyr Met Ala Gly Arg Val Ala Thr Ser Thr Ser Gly Cys His Cys Trp  
20 25 30  
Met Ser Arg Arg Asp Leu Thr Pro Leu Pro His Pro Ser Glu Pro Gly  
35 40 45  
Val Leu Asp Cys Leu Gly Pro Cys His Leu Leu Pro Leu Leu Ser Pro  
50 55 60  
Gly Ser Pro Cys Trp Val Leu Gly Leu His Phe Ser Leu His Pro Pro  
65 70 75 80  
Ser Ala Ala Ser Ala Ser His Ala Leu Thr Ile Thr Ser Leu Pro Pro  
85 90 95  
Gly Leu Leu Pro Phe Val Gly Val Glu Leu Thr Ala His Pro Gln Ala  
100 105 110  
Leu Ile Gly Arg Gly Phe Pro Ser Gly Met Ala Ala Ala Gly Arg His  
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Leu Cys Phe Leu  
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ctgctggatg tcaagacgtg a 81

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<400> 56  
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 Lys Arg Met Pro Leu Leu Asp Val Lys Thr  
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 ctgcttcccc tgctgtcacc tggctcccc tgctgggtgc tgggtctcca tttctcctc 180  
 caccacccct cagcagcatc tgcttcccat gccctcacca tcacctcact gccccaggc 240  
 cttctgccct ttgtgggtgt tgagctcacc gccacccac aggcactcat aggaagaggc 300  
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 35 40 45  
 Ser Pro Cys Trp Val Leu Gly Leu His Phe Ser Leu His Pro Pro Ser  
 50 55 60  
 Ala Ala Ser Ala Ser His Ala Leu Thr Ile Thr Ser Leu Pro Pro Gly  
 65 70 75 80  
 Leu Leu Pro Phe Val Gly Val Glu Leu Thr Ala His Pro Gln Ala Leu  
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 Gly Ser Pro Cys Trp Val Leu Gly Leu His Phe Ser Leu His Pro Pro  
 35 40 45  
 Ser Ala Ala Ser Ala Ser His Ala Leu Thr Ile Thr Ser Leu Pro Pro  
 50 55 60  
 Gly Leu Leu Pro Phe Val Gly Val Glu Leu Thr Ala His Pro Gln Ala  
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<400> 64  
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<210> 69  
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<400> 69  
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accatccata	cgattcagtt	gattcagcac	aaccgacgtc	ttcgcaacct	tattgccaca		240
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ccctcgtgcc	ctgggtcacc	tctctccct	gatgacctcc	tgcctttaga	ttgtaagaat		360
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gactcagata	gcagctatgg	ttcccactcc	actgacagcc	tcatggggtc	ctcccctgtt		1200
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<210> 77
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			20					25					30		
His	Asp	Pro	Pro	Leu	His	Gln	Pro	Ser	Ala	Asn	Lys	Pro	Lys	Pro	Pro
		35					40					45			
Thr	Met	Leu	Asp	Ile	Pro	Ser	Glu	Pro	Cys	Ser	Leu	Thr	Ile	His	Thr
	50					55					60				
Ile	Gln	Leu	Ile	Gln	His	Asn	Arg	Arg	Leu	Arg	Asn	Leu	Ile	Ala	Thr
65					70					75					80
Ala	Gln	Ala	Gln	Asn	Gln	Gln	Gln	Thr	Glu	Gly	Val	Lys	Thr	Glu	Glu
				85					90					95	
Ser	Glu	Pro	Leu	Pro	Ser	Cys	Pro	Gly	Ser	Pro	Pro	Leu	Pro	Asp	Asp
			100					105					110		
Leu	Leu	Pro	Leu	Asp	Cys	Lys	Asn	Pro	Asn	Ala	Pro	Phe	Gln	Ile	Arg
		115					120					125			
His	Ser	Asp	Pro	Glu	Ser	Asp	Phe	Tyr	Arg	Gly	Lys	Gly	Glu	Pro	Val
	130					135					140				
Thr	Glu	Leu	Ser	Trp	His	Ser	Cys	Arg	Gln	Leu	Leu	Tyr	Gln	Ala	Val
145					150					155					160
Ala	Thr	Ile	Leu	Ala	His	Ala	Gly	Phe	Asp	Cys	Ala	Asn	Glu	Ser	Val
				165					170					175	
Leu	Glu	Thr	Leu	Thr	Asp	Val	Ala	His	Glu	Tyr	Cys	Leu	Lys	Phe	Thr
			180					185					190		
Lys	Leu	Leu	Arg	Phe	Ala	Val	Asp	Arg	Glu	Ala	Arg	Leu	Gly	Gln	Thr
		195					200					205			
Pro	Phe	Pro	Asp	Val	Met	Glu	Gln	Val	Phe	His	Glu	Val	Gly	Ile	Gly
	210					215					220				
Ser	Val	Leu	Ser	Leu	Gln	Lys	Phe	Trp	Gln	His	Arg	Ile	Lys	Asp	Tyr
225					230					235					240
His	Ser	Tyr	Met	Leu	Gln	Ile	Ser	Lys	Gln	Leu	Ser	Glu	Glu	Tyr	Glu
			245						250					255	
Arg	Ile	Val	Asn	Pro	Glu	Lys	Ala	Thr	Glu	Asp	Ala	Lys	Pro	Val	Lys
			260					265					270		
Ile	Lys	Glu	Glu	Pro	Val	Ser	Asp	Ile	Thr	Phe	Pro	Val	Ser	Glu	Glu
		275					280					285			
Leu	Glu	Ala	Asp	Leu	Ala	Ser	Gly	Asp	Gln	Ser	Leu	Pro	Met	Gly	Val
		290				295					300				
Leu	Gly	Ala	Gln	Ser	Glu	Arg	Phe	Pro	Ser	Asn	Leu	Glu	Val	Glu	Ala
305					310					315					320
Ser	Pro	Gln	Ala	Ser	Ser	Ala	Glu	Val	Asn	Ala	Ser	Pro	Leu	Trp	Asn
				325					330					335	
Leu	Ala	His	Val	Lys	Met	Glu	Pro	Gln	Glu	Ser	Glu	Glu	Gly	Asn	Val
			340					345					350		
Ser	Gly	His	Gly	Val	Leu	Gly	Ser	Asp	Val	Phe	Glu	Glu	Pro	Met	Ser
		355					360					365			
Gly	Met	Ser	Glu	Ala	Gly	Ile									

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<210> 78
<211> 105
<212> DNA
<213> Homo sapiens
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<400> 78  
 atgacccacc cctgcaccaa ccctcagcca acaagccgaa gccccccact atgctggaca 60  
 tcccctcaga gccatgtagt ctcaccatcc atacgattca gttga 105

<210> 79  
 <211> 34  
 <212> PRT  
 <213> Homo sapiens

<400> 79  
 Met Thr His Pro Cys Thr Asn Pro Gln Pro Thr Ser Arg Ser Pro Pro  
 1 5 10 15  
 Leu Cys Trp Thr Ser Pro Gln Ser His Val Val Ser Pro Ser Ile Arg  
 20 25 30  
 Phe Ser

<210> 80  
 <211> 1092  
 <212> DNA  
 <213> Homo sapiens

<400> 80  
 atgctggaca tcccctcaga gccatgtagt ctcaccatcc atacgattca gttgattcag 60  
 cacaaccgac gtcttcgcaa ccttattgcc acagctcagg cccagaatca gcagcagaca 120  
 gaaggtgtaa aaactgaaga gagtgaacct cttccctcgt gccctgggtc acctcctctc 180  
 cctgatgacc tcctgccttt agattgtaag aatcccaatg caccattcca gatccggcac 240  
 agtgacccag agagtgactt ttatcgtggg aaaggggaac ctgtgactga actcagctgg 300  
 cactcctgtc ggcagctcct ctaccaggca gtggccacaa tcctggccca cgcgggcttt 360  
 gactgtgcta atgagagtgt cctggagacc ctaactgatg tggcacatga gtattgcctt 420  
 aagtttacca agttgctgcg ttttgctgtg gaccgggagg cccggctggg acagactcct 480  
 tttcctgatg tgatggagca ggtattccat gaagtgggta ttggcagtgt gctctccctc 540  
 cagaagttct ggcagcaccg catcaaggac tatcacagtt acatgctaca gattagtaag 600  
 caactctctg aagaatatga aaggattgtc aatcctgaga aggccacaga ggacgctaaa 660  
 cctgtgaaga tcaaggagga acctgtgagc gacatcactt ttctgtcag tgaggagctg 720  
 gaggtgacc ttgcttctgg agaccagtca ctgcctatgg gagtgcttgg ggctcagagc 780  
 gaacgcttcc catctaacct ggaggttgaa gcttcaccac aggcttcaag tgcagaggta 840  
 aatgcttctc ctcttttgaa tctggcccat gtgaaaatgg agcctcaaga aagtgaagaa 900  
 ggcaatgtct ctgggcatgg tgtgctgggc agtgatgtct tcgaggagcc tatgtcaggc 960  
 atgagtgaag ctgggattcc tcagagccct gatgactcag atagcagcta tggttcccac 1020  
 tccactgaca gcctcatggg gtccctccct gttttcaacc agcgctgcaa gaagaggatg 1080  
 aggaaaatat aa 1092

<210> 81  
 <211> 363  
 <212> PRT  
 <213> Homo sapiens

<400> 81  
 Met Leu Asp Ile Pro Ser Glu Pro Cys Ser Leu Thr Ile His Thr Ile  
 1 5 10 15  
 Gln Leu Ile Gln His Asn Arg Arg Leu Arg Asn Leu Ile Ala Thr Ala  
 20 25 30  
 Gln Ala Gln Asn Gln Gln Gln Thr Glu Gly Val Lys Thr Glu Glu Ser  
 35 40 45

Glu	Pro	Leu	Pro	Ser	Cys	Pro	Gly	Ser	Pro	Pro	Leu	Pro	Asp	Asp	Leu
50						55					60				
Leu	Pro	Leu	Asp	Cys	Lys	Asn	Pro	Asn	Ala	Pro	Phe	Gln	Ile	Arg	His
65				70					75						80
Ser	Asp	Pro	Glu	Ser	Asp	Phe	Tyr	Arg	Gly	Lys	Gly	Glu	Pro	Val	Thr
				85					90					95	
Glu	Leu	Ser	Trp	His	Ser	Cys	Arg	Gln	Leu	Leu	Tyr	Gln	Ala	Val	Ala
			100					105					110		
Thr	Ile	Leu	Ala	His	Ala	Gly	Phe	Asp	Cys	Ala	Asn	Glu	Ser	Val	Leu
		115				120						125			
Glu	Thr	Leu	Thr	Asp	Val	Ala	His	Glu	Tyr	Cys	Leu	Lys	Phe	Thr	Lys
		130				135					140				
Leu	Leu	Arg	Phe	Ala	Val	Asp	Arg	Glu	Ala	Arg	Leu	Gly	Gln	Thr	Pro
145					150					155					160
Phe	Pro	Asp	Val	Met	Glu	Gln	Val	Phe	His	Glu	Val	Gly	Ile	Gly	Ser
				165					170					175	
Val	Leu	Ser	Leu	Gln	Lys	Phe	Trp	Gln	His	Arg	Ile	Lys	Asp	Tyr	His
			180					185					190		
Ser	Tyr	Met	Leu	Gln	Ile	Ser	Lys	Gln	Leu	Ser	Glu	Glu	Tyr	Glu	Arg
		195					200					205			
Ile	Val	Asn	Pro	Glu	Lys	Ala	Thr	Glu	Asp	Ala	Lys	Pro	Val	Lys	Ile
		210				215					220				
Lys	Glu	Glu	Pro	Val	Ser	Asp	Ile	Thr	Phe	Pro	Val	Ser	Glu	Glu	Leu
225					230					235					240
Glu	Ala	Asp	Leu	Ala	Ser	Gly	Asp	Gln	Ser	Leu	Pro	Met	Gly	Val	Leu
				245					250					255	
Gly	Ala	Gln	Ser	Glu	Arg	Phe	Pro	Ser	Asn	Leu	Glu	Val	Glu	Ala	Ser
			260					265					270		
Pro	Gln	Ala	Ser	Ser	Ala	Glu	Val	Asn	Ala	Ser	Pro	Leu	Trp	Asn	Leu
		275					280					285			
Ala	His	Val	Lys	Met	Glu	Pro	Gln	Glu	Ser	Glu	Glu	Gly	Asn	Val	Ser
		290				295					300				
Gly	His	Gly	Val	Leu	Gly	Ser	Asp	Val	Phe	Glu	Glu	Pro	Met	Ser	Gly
305					310					315					320
Met	Ser	Glu	Ala	Gly	Ile	Pro	Gln	Ser	Pro	Asp	Asp	Ser	Asp	Ser	Ser
				325					330					335	
Tyr	Gly	Ser	His	Ser	Thr	Asp	Ser	Leu	Met	Gly	Ser	Ser	Pro	Val	Phe
			340					345					350		
Asn	Gln	Arg	Cys	Lys	Lys	Arg	Met	Arg	Lys	Ile					
		355					360								

<210> 82  
 <211> 18  
 <212> DNA  
 <213> Homo sapiens

<400> 82  
 atgacctcct gcctttag

18

<210> 83  
 <211> 5  
 <212> PRT  
 <213> Homo sapiens

<400> 83

Met Thr Ser Cys Leu  
1 5

<210> 84  
<211> 69  
<212> DNA  
<213> Homo sapiens

<400> 84  
atgcaccatt ccagatccgg cacagtgacc cagagagtga cttttatcgt gggaaagggg 60  
aacctgtga 69

<210> 85  
<211> 22  
<212> PRT  
<213> Homo sapiens

<400> 85  
Met His His Ser Arg Ser Gly Thr Val Thr Gln Arg Val Thr Phe Ile  
1 5 10 15  
Val Gly Lys Gly Asn Leu  
20

<210> 86  
<211> 24  
<212> DNA  
<213> Homo sapiens

<400> 86  
atgagagtgt cctggagacc ctaa 24

<210> 87  
<211> 7  
<212> PRT  
<213> Homo sapiens

<400> 87  
Met Arg Val Ser Trp Arg Pro  
1 5

<210> 88  
<211> 96  
<212> DNA  
<213> Homo sapiens

<400> 88  
atgtggcaca tgagtattgc cttaagttaa ccaagttgct gcgttttgct gtggaccggg 60  
aggccccggct gggacagact ccttttcctg atgtga 96

<210> 89  
<211> 31  
<212> PRT  
<213> Homo sapiens

<400> 89

Met Trp His Met Ser Ile Ala Leu Ser Leu Pro Ser Cys Cys Val Leu  
 1 5 10 15  
 Leu Trp Thr Gly Arg Pro Gly Trp Asp Arg Leu Leu Phe Leu Met  
 20 25 30

<210> 90  
 <211> 87  
 <212> DNA  
 <213> Homo sapiens

<400> 90  
 atgagtattg ccttaagttt accaagttgc tgcgttttgc tgtggaccgg gagggccggc 60  
 tgggacagac tccttttcct gatgtga 87

<210> 91  
 <211> 28  
 <212> PRT  
 <213> Homo sapiens

<400> 91  
 Met Ser Ile Ala Leu Ser Leu Pro Ser Cys Cys Val Leu Leu Trp Thr  
 1 5 10 15  
 Gly Arg Pro Gly Trp Asp Arg Leu Leu Phe Leu Met  
 20 25

<210> 92  
 <211> 600  
 <212> DNA  
 <213> Homo sapiens

<400> 92  
 atggagcagg tattccatga agtgggtatt ggcagtgtgc tctccctcca gaagttctgg 60  
 cagcaccgca tcaaggacta tcacagttac atgctacaga ttagtaagca actctctgaa 120  
 gaatatgaaa ggattgtcaa tcctgagaag gccacagagg acgctaaacc tgtgaagatc 180  
 aaggaggaac ctgtgagcga catcactttt cctgtcagtg aggagctgga ggctgacctt 240  
 gcttctggag accagtcact gcctatggga gtgcttgggg ctcagagcga acgcttccca 300  
 tctaacctgg aggttgaagc ttcaccacag gcttcaagtg cagaggtaaa tgcttctcct 360  
 ctttggaatc tggcccatgt gaaaatggag cctcaagaaa gtgaagaagg caatgtctct 420  
 gggcatggtg tgctgggcag tgatgtcttc gaggagccta tgtcaggcat gagtgaagct 480  
 gggattcctc agagccctga tgactcagat agcagctatg gttcccactc cactgacagc 540  
 ctcatggggt cctcccctgt tttcaaccag cgctgcaaga agaggatgag gaaaatataa 600

<210> 93  
 <211> 199  
 <212> PRT  
 <213> Homo sapiens

<400> 93  
 Met Glu Gln Val Phe His Glu Val Gly Ile Gly Ser Val Leu Ser Leu  
 1 5 10 15  
 Gln Lys Phe Trp Gln His Arg Ile Lys Asp Tyr His Ser Tyr Met Leu  
 20 25 30  
 Gln Ile Ser Lys Gln Leu Ser Glu Tyr Glu Arg Ile Val Asn Pro  
 35 40 45  
 Glu Lys Ala Thr Glu Asp Ala Lys Pro Val Lys Ile Lys Glu Glu Pro







<213> Homo sapiens

<400> 100

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atgggagtgctc ttgggggctca gagcgaacgc ttcccatcta acctggaggt tgaagcttca      60
ccacaggctt caagtgcaga ggtaaagtct tctcctcttt ggaatctggc ccatgtgaaa      120
atggagcctc aagaaagtga agaaggcaat gtctctgggc atggtgtgct gggcagtgat      180
gtcttcgagg agcctatgtc aggcattgagt gaagctggga ttcttcagag ccctgatgac      240
tcagatagca gctatggttc ccactccact gacagcctca tggggtcctc ccctgttttc      300
aaccagcgct gcaagaagag gatgaggaaa atataa      336
```

<210> 101

<211> 111

<212> PRT

<213> Homo sapiens

<400> 101

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Met Gly Val Leu Gly Ala Gln Ser Glu Arg Phe Pro Ser Asn Leu Glu
 1           5           10           15
Val Glu Ala Ser Pro Gln Ala Ser Ser Ala Glu Val Asn Ala Ser Pro
      20           25           30
Leu Trp Asn Leu Ala His Val Lys Met Glu Pro Gln Glu Ser Glu Glu
      35           40           45
Gly Asn Val Ser Gly His Gly Val Leu Gly Ser Asp Val Phe Glu Glu
      50           55           60
Pro Met Ser Gly Met Ser Glu Ala Gly Ile Pro Gln Ser Pro Asp Asp
      65           70           75           80
Ser Asp Ser Ser Tyr Gly Ser His Ser Thr Asp Ser Leu Met Gly Ser
      85           90           95
Ser Pro Val Phe Asn Gln Arg Cys Lys Lys Arg Met Arg Lys Ile
      100           105           110
```

<210> 102

<211> 33

<212> DNA

<213> Homo sapiens

<400> 102

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atgcttctcc tctttggaat ctggcccatg tga      33
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<210> 103

<211> 10

<212> PRT

<213> Homo sapiens

<400> 103

```
Met Leu Leu Leu Phe Gly Ile Trp Pro Met
 1           5           10
```

<210> 104

<211> 216

<212> DNA

<213> Homo sapiens

<400> 104

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atggagcctc aagaaagtga agaaggcaat gtctctgggc atggtgtgct gggcagtgat      60
```

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gtcttcgagg agcctatgtc aggcattgagt gaagctggga ttcctcagag ccctgatgac 120
tcagatagca gctatgggtc ccaactccact gacagcctca tgggggtcctc ccctgttttc 180
aaccagcgct gcaagaagag gatgaggaaa atataa 216
```

<210> 105  
 <211> 71  
 <212> PRT  
 <213> Homo sapiens

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<400> 105
Met Glu Pro Gln Glu Ser Glu Glu Gly Asn Val Ser Gly His Gly Val
 1          5          10          15
Leu Gly Ser Asp Val Phe Glu Glu Pro Met Ser Gly Met Ser Glu Ala
 20          25          30
Gly Ile Pro Gln Ser Pro Asp Asp Ser Asp Ser Ser Tyr Gly Ser His
 35          40          45
Ser Thr Asp Ser Leu Met Gly Ser Ser Pro Val Phe Asn Gln Arg Cys
 50          55          60
Lys Lys Arg Met Arg Lys Ile
65          70
```

<210> 106  
 <211> 60  
 <212> DNA  
 <213> Homo sapiens

```
<400> 106
atgtctctgg gcatggtgtg ctgggcagtg atgtcttcga ggagcctatg tcaggcatga 60
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<210> 107  
 <211> 19  
 <212> PRT  
 <213> Homo sapiens

```
<400> 107
Met Ser Leu Gly Met Val Cys Trp Ala Val Met Ser Ser Arg Ser Leu
 1          5          10          15
Cys Gln Ala
```

<210> 108  
 <211> 48  
 <212> DNA  
 <213> Homo sapiens

```
<400> 108
atggtgtgct gggcagtgat gtcttcgagg agcctatgtc aggcattgac 48
```

<210> 109  
 <211> 15  
 <212> PRT  
 <213> Homo sapiens

```
<400> 109
Met Val Cys Trp Ala Val Met Ser Ser Arg Ser Leu Cys Gln Ala
```



<213> Homo sapiens

<400> 115

Met Ser Glu Ala Gly Ile Pro Gln Ser Pro Asp Asp Ser Asp Ser Ser  
1 5 10 15  
Tyr Gly Ser His Ser Thr Asp Ser Leu Met Gly Ser Ser Pro Val Phe  
20 25 30  
Asn Gln Arg Cys Lys Lys Arg Met Arg Lys Ile  
35 40

<210> 116

<211> 90

<212> DNA

<213> Homo sapiens

<400> 116

atgactcaga tagcagctat ggttcccact ccactgacag cctcatgggg tcttccctg 60  
ttttcaacca gcgctgcaag aagaggatga 90

<210> 117

<211> 29

<212> PRT

<213> Homo sapiens

<400> 117

Met Thr Gln Ile Ala Ala Met Val Pro Thr Pro Leu Thr Ala Ser Trp  
1 5 10 15  
Gly Pro Pro Leu Phe Ser Thr Ser Ala Ala Arg Arg Gly  
20 25

<210> 118

<211> 72

<212> DNA

<213> Homo sapiens

<400> 118

atgggttccca ctccactgac agcctcatgg ggtcctcccc tgttttcaac cagcgctgca 60  
agaagaggat ga 72

<210> 119

<211> 23

<212> PRT

<213> Homo sapiens

<400> 119

Met Val Pro Thr Pro Leu Thr Ala Ser Trp Gly Pro Pro Leu Phe Ser  
1 5 10 15  
Thr Ser Ala Ala Arg Arg Gly  
20

<210> 120

<211> 57

<212> DNA

<213> Homo sapiens





Glu Val Arg Ser Ser Arg Pro Ala  
20

<210> 132  
<211> 39  
<212> DNA  
<213> Homo sapiens

<400> 132  
atggaaaaaa ccccatctct actaaaaata caaaattag

39

<210> 133  
<211> 12  
<212> PRT  
<213> Homo sapiens

<400> 133  
Met Glu Lys Thr Pro Ser Leu Leu Lys Ile Gln Asn  
1 5 10

<210> 134  
<211> 33  
<212> DNA  
<213> Homo sapiens

<400> 134  
atgcctgtaa tcccagctac tcaggaaggc tga

33

<210> 135  
<211> 10  
<212> PRT  
<213> Homo sapiens

<400> 135  
Met Pro Val Ile Pro Ala Thr Gln Glu Gly  
1 5 10

<210> 136  
<211> 542  
<212> DNA  
<213> Homo sapiens

<400> 136  
tcgacccacg cgtccgggac aatagtgtag gttatggatg gaggtgtcgg tactaaattg 60  
aataacgagt aaataatctt acttgggtag agatggcctt tgccaacaaa gtgaactgtt 120  
ttggttggtt taaactcatg aagtatgggt tcagtggaaa tgtttggaac tctgaaggat 180  
ttagacaagg ttttgaaaag gataatcatg ggtagaagg aagtgtttga aagtcacttt 240  
gaaagttagt tttgggccag cacggtagct cacccttgta atcccagcac tttgggaggc 300  
tgaggtgggt agattacttg agcccaggaa ttcaagacca gcctgggcaa catggtgaaa 360  
ccctgtttct ataaaaaata atctgggctt tgtagcatat gcctgtgggc ccagctactg 420  
aggaggctga ggtgggagga ttgcttgagc ccaggaggca gaggttgagc tgagccaagg 480  
tcagtcact gcactctagc ctgggcaaca gagtaagaca aaaaaaaaaa aaaagggcgg 540  
cc 542

<210> 137





[illegible]60  
93

93



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aagaagtctt gaatagctct ttactgtctt acttgggggtt gataagattt gagtgtttgc 540
aatttttttac taaatgtagc tccaaagtct taaatggctt gtttgttctt aaactgttaa 600
ttgatgaaac tgtgcataag ttacaatgt actaacttat tttgcttatt atatatagtg 660
ttttattgga aattgtaacc acacacttca gcatgatgaa aataaagatt agtgtttcca 720
tttaaataaa tgttttatcc tcccataaaa aaaaaaaaaa aaagggcggc c 771

```

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<210> 154
<211> 108
<212> DNA
<213> Homo sapiens

```

```

<400> 154
atgtgtccca tgtgggttgt gccaggtaga gaaacaggaa gtcaatcatc tgtgacagtc 60
tctattctgt cgttttgctc cttggtattt gatttgact atatttag 108

```

```

<210> 155
<211> 35
<212> PRT
<213> Homo sapiens

```

```

<400> 155
Met Cys Pro Met Trp Val Val Pro Gly Arg Glu Thr Gly Ser Gln Ser
  1             5             10             15
Ser Val Thr Val Ser Ile Leu Ser Phe Cys Ser Leu Val Phe Asp Leu
      20             25             30
His Tyr Ile
      35

```

```

<210> 156
<211> 99
<212> DNA
<213> Homo sapiens

```

```

<400> 156
atgtgggttg tgccaggtag agaaacagga agtcaatcat ctgtgacagt ctctattctg 60
tcgttttgct ccttggtatt tgatttgcac tatatttag 99

```

```

<210> 157
<211> 32
<212> PRT
<213> Homo sapiens

```

```

<400> 157
Met Trp Val Val Pro Gly Arg Glu Thr Gly Ser Gln Ser Ser Val Thr
  1             5             10             15
Val Ser Ile Leu Ser Phe Cys Ser Leu Val Phe Asp Leu His Tyr Ile
      20             25             30

```

```

<210> 158
<211> 21
<212> DNA
<213> Homo sapiens

```

```

<400> 158
atggagacct ggttccagta a 21

```

<210> 159  
 <211> 6  
 <212> PRT  
 <213> Homo sapiens

<400> 159  
 Met Glu Thr Trp Phe Gln  
 1 5

<210> 160  
 <211> 75  
 <212> DNA  
 <213> Homo sapiens

<400> 160  
 atgtcccacc agtgggggtat agaaagcatg ctcattgaccc tgccgtgtcg tctgaggtac 60  
 ccgttcttat cctag 75

<210> 161  
 <211> 24  
 <212> PRT  
 <213> Homo sapiens

<400> 161  
 Met Ser His Gln Trp Gly Ile Glu Ser Met Leu Met Thr Leu Pro Cys  
 1 5 10 15  
 Arg Leu Arg Tyr Pro Phe Leu Ser  
 20

<210> 162  
 <211> 48  
 <212> DNA  
 <213> Homo sapiens

<400> 162  
 atgtctcatga ccctgccgtg tcgtctgagg taccggttct taccctag 48

<210> 163  
 <211> 15  
 <212> PRT  
 <213> Homo sapiens

<400> 163  
 Met Leu Met Thr Leu Pro Cys Arg Leu Arg Tyr Pro Phe Leu Ser  
 1 5 10 15

<210> 164  
 <211> 42  
 <212> DNA  
 <213> Homo sapiens

<400> 164  
 atgaccctgc cgtgtcgtct gaggtaccgc ttcttattcct ag 42

<210> 165

**L** **E** **A** **R** **E** **S**

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<210> 166
<211> 99
<212> DNA
<213> Homo sapiens
```

```
<210> 167
<211> 32
<212> PRT
<213> Homo sapiens
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```
<210> 168
<211> 63
<212> DNA
<213> Homo sapiens
```

```
<210> 169
<211> 20
<212> PRT
<213> Homo sapiens
```

```
<210> 170
<211> 39
<212> DNA
<213> Homo sapiens
```

37

Instrument	Part	Key	Tempo	Time
Violin I	First Violin	D Major	Allegro	1:00
Violin II	Second Violin	D Major	Allegro	1:00
Viola	Viola	D Major	Allegro	1:00
Cello	Cello	D Major	Allegro	1:00
Double Bass	Double Bass	D Major	Allegro	1:00
Piano	Piano	D Major	Allegro	1:00
Conductor	Conductor	D Major	Allegro	1:00

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<210> 172
<211> 177
<212> DNA
<213> Homo sapiens
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<210> 173
<211> 58
<212> PRT
<213> Homo sapiens
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<210> 174
<211> 27
<212> DNA
<213> Homo sapiens
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<210> 175
<211> 8
<212> PRT
<213> Homo sapiens
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<210> 176
<211> 75
<212> DNA
<213> Homo sapiens
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atgtttttatc ctcccataaa aaaaaaaaaa aaaagggcgg cc

42

<210> 183

<211> 14

<212> PRT

<213> Homo sapiens

<400> 183

Met Phe Tyr Pro Pro Ile Lys Lys Lys Lys Lys Arg Ala Ala

1

5

10

<210> 184

<211> 1669

<212> DNA

<213> Homo sapiens

<400> 184

tcgacccacg	cgtccgcagg	cagtgcactgc	cttcggccttt	ttttctgctg	actaagatct	60
cctatagaga	gctacaacaa	tgcccaaaag	aaagccaaag	agaagatctg	ccaggttgct	120
tgctatgctt	gtgccagtta	caccagaggt	gaagcctaaa	agaacatcaa	gttcaaggaa	180
aatgaagaca	aaaagtgata	tgatggaaga	aaacatagat	acaagtgcc	aagcagttgc	240
tgaaaaccaag	caagaagcag	ttgttggaaga	agactacaat	gaaaatgcta	aaaatggaga	300
agccaaaatt	acagaggcac	cagcttctga	aaaagaaatt	gtggaagtaa	aagaagaaaa	360
tattgaagat	gccacagaaa	agggaggaga	aaagaaagaa	gcagtggcag	cagaagtaaa	420
aaatgaagaa	gaagatcaga	aagaagatga	agaagatcaa	aacgaagaga	aaggggaagc	480
tggaagaaag	gacaaagatg	aaaaagggga	agaagatgga	aaagaggata	aaaatggaaa	540
tgagaaagga	gaagatgcaa	aagagaaaga	agatggaaaa	aaaggtgaag	acggaaaagg	600
aaatggagaa	gatggaaaag	agaaaggaga	agatgaaaaa	gaggaagaag	acagaaaaga	660
aacaggagtt	ggaaaagaga	atgaagatgg	aaaagagaag	ggagataaaa	aagaggggaa	720
agatgtaaaa	gtcaaagaag	atgaaaaaga	gagagaagat	ggaaaagaag	atgaaggtgg	780
aaatgaggaa	gaagctggaa	aagagaaaga	agatttaaaa	gaagaggaag	aaggaaaaaga	840
ggaagatgag	atcaaagaag	atgatggaaa	aaaagaggag	ccacagagta	ttgtttaaaa	900
ctgccctatg	tagtttcata	atttggtaac	atgtaccttc	atgttgtaaa	gttaatatag	960
ataaatatatt	ttatcaaaaa	ttttataaac	acagcctttc	tttagcattg	atttaatttc	1020
agaacatctt	catattgatt	attagccata	aagtttctaa	catgaaacat	ttatctataa	1080
attttgtgat	tatagtagtg	gaatacatag	aaaaaaatat	gctttcaact	ttgtgagtga	1140
atttcgtggt	gtgtaagtta	tatgtcaaat	ctttgaattt	taattttact	cctttttatac	1200
atgtgataat	ttcataaagt	gagggatccc	aaaaaaagag	tttcatccca	acattcttgt	1260
tctgcagggt	gctttttataa	agaaggtgaa	ctatttttcat	gtaatgttaa	gagttaaact	1320
tatctttccc	aaatataact	ttattattag	cttgggaaaa	atgaaattgt	attcccattt	1380
ttaaaataaaa	tacaaatggt	tatttcagaa	gggcagtttt	gattatatgt	gaatacacaa	1440
attttactgg	atttatctta	ataaaaagac	tctgacgatg	attgtgtttt	gttatatctt	1500
caaaaatata	gctagtgaag	tattgtgctt	aatttttttc	tattgtgtta	ttcatgaaaa	1560
tatttaatat	tactgacat	aaaattaata	taaagtaaaa	ttcaccattt	taattataat	1620
aaaaataaag	tatataattc	aaaaaaaaaa	aaaaaaaaaa	agggcggcc		1669

<210> 185

<211> 819

<212> DNA

<213> Homo sapiens

<400> 185

atgccccaaa	gaaagccaaa	gagaagatct	gccaggttgt	ctgctatgct	tgtgccagtt	60
acaccagagg	tgaagcctaa	aagaacatca	agttcaagga	aatgaagac	aaaaagtgat	120
atgatggaag	aaaacataga	tacaagtgcc	caagcagttg	ctgaaaccaa	gcaagaagca	180



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gttgttgaag aagactacaa tgaaaatgct aaaaatggag aagccaaaat tacagaggca 240
ccagcttctg aaaaagaaat tgtggaagta aaagaagaaa atattgaaga tgccacagaa 300
aagggaggag aaaagaaaga agcagtggca gcagaagtaa aaaatgaaga agaagatcag 360
aaagaagatg aagaagatca aaacgaagag aaaggggaag ctggaaaaga agacaaagat 420
gaaaaagggg aagaagatgg aaaagaggat aaaaatggaa atgagaaagg agaagatgca 480
aaagagaaaag aagatggaaa aaaaggtgaa gacggaaaag gaaatggaga agatggaaaa 540
gagaaaggag aagatgaaaa agaggaagaa gacagaaaag aaacaggagt tggaaaagag 600
aatgaagatg gaaaagagaa gggagataaa aaagagggga aagatgtaaa agtcaaagaa 660
gatgaaaaag agagagaaga tggaaaagaa gatgaaggtg gaaatgagga agaagctgga 720
aaagagaaaag aagatttaaa agaagaggaa gaaggaaaag aggaagatga gatcaaagaa 780
gatgatggaa aaaaagagga gccacagagt attgtttaa 819

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<210> 186  
 <211> 272  
 <212> PRT  
 <213> Homo sapiens

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<400> 186
Met Pro Lys Arg Lys Pro Lys Arg Arg Ser Ala Arg Leu Ser Ala Met
  1          5          10          15
Leu Val Pro Val Thr Pro Glu Val Lys Pro Lys Arg Thr Ser Ser Ser
          20          25          30
Arg Lys Met Lys Thr Lys Ser Asp Met Met Glu Glu Asn Ile Asp Thr
          35          40          45
Ser Ala Gln Ala Val Ala Glu Thr Lys Gln Glu Ala Val Val Glu Glu
          50          55          60
Asp Tyr Asn Glu Asn Ala Lys Asn Gly Glu Ala Lys Ile Thr Glu Ala
          65          70          75          80
Pro Ala Ser Glu Lys Glu Ile Val Glu Val Lys Glu Glu Asn Ile Glu
          85          90          95
Asp Ala Thr Glu Lys Gly Gly Glu Lys Lys Glu Ala Val Ala Ala Glu
          100          105          110
Val Lys Asn Glu Glu Glu Asp Gln Lys Glu Asp Glu Glu Asp Gln Asn
          115          120          125
Glu Glu Lys Gly Glu Ala Gly Lys Glu Asp Lys Asp Glu Lys Gly Glu
          130          135          140
Glu Asp Gly Lys Glu Asp Lys Asn Gly Asn Glu Lys Gly Glu Asp Ala
          145          150          155          160
Lys Glu Lys Glu Asp Gly Lys Lys Gly Glu Asp Gly Lys Gly Asn Gly
          165          170          175
Glu Asp Gly Lys Glu Lys Gly Glu Asp Glu Lys Glu Glu Glu Asp Arg
          180          185          190
Lys Glu Thr Gly Val Gly Lys Glu Asn Glu Asp Gly Lys Glu Lys Gly
          195          200          205
Asp Lys Lys Glu Gly Lys Asp Val Lys Val Lys Glu Asp Glu Lys Glu
          210          215          220
Arg Glu Asp Gly Lys Glu Asp Glu Gly Gly Asn Glu Glu Glu Ala Gly
          225          230          235          240
Lys Glu Lys Glu Asp Leu Lys Glu Glu Glu Glu Gly Lys Glu Glu Asp
          245          250          255
Glu Ile Lys Glu Asp Asp Gly Lys Lys Glu Glu Pro Gln Ser Ile Val
          260          265          270

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<210> 187  
 <211> 774

<212> DNA  
<213> Homo sapiens

<400> 187  
atgcttgtgc cagttacacc agaggtgaag cctaaaagaa catcaagttc aaggaaaatg 60  
aagacaaaaa gtgatatgat ggaagaaaac atagatacaa gtgccaagc agtttgctgaa 120  
accaagcaag aagcagttgt tgaagaagac tacaatgaaa atgctaaaaa tggagaagcc 180  
aaaattacag aggcaccagc ttctgaaaaa gaaatttgtg aagtaaaaga agaaaaatatt 240  
gaagatgcc aagaaaagg aggagaaaag aaagaagcag tggcagcaga agtaaaaaat 300  
gaagaagaag atcagaaaaga agatgaagaa gatcaaacg aagagaaaag ggaagctgga 360  
aaagaagaca aagatgaaaa aggggaagaa gatggaaaag aggataaaaa tggaaatgag 420  
aaaggagaag atgcaaaaaga gaaagaagat ggaaaaaaag gtgaagacgg aaaaggaaat 480  
ggagaagatg gaaaagagaa aggagaagat gaaaaagagg aagaagacag aaaagaaaca 540  
ggagttggaa aagagaatga agatggaaaa gagaaggagg ataaaaaaga ggggaaagat 600  
gtaaaagtca aagaagatga aaaagagaga gaagatggaa aagaagatga aggtggaaat 660  
gaggaagaag ctggaaaaga gaaagaagat taaaagaag aggaagaagg aaaagaggaa 720  
gatgagatca aagaagatga tggaaaaaaa gaggagccac agagtattgt ttaa 774

<210> 188  
<211> 257  
<212> PRT  
<213> Homo sapiens

<400> 188  
Met Leu Val Pro Val Thr Pro Glu Val Lys Pro Lys Arg Thr Ser Ser  
1 5 10 15  
Ser Arg Lys Met Lys Thr Lys Ser Asp Met Met Glu Glu Asn Ile Asp  
20 25 30  
Thr Ser Ala Gln Ala Val Ala Glu Thr Lys Gln Glu Ala Val Val Glu  
35 40 45  
Glu Asp Tyr Asn Glu Asn Ala Lys Asn Gly Glu Ala Lys Ile Thr Glu  
50 55 60  
Ala Pro Ala Ser Glu Lys Glu Ile Val Glu Val Lys Glu Glu Asn Ile  
65 70 75 80  
Glu Asp Ala Thr Glu Lys Gly Gly Glu Lys Lys Glu Ala Val Ala Ala  
85 90 95  
Glu Val Lys Asn Glu Glu Glu Asp Gln Lys Glu Asp Glu Glu Asp Gln  
100 105 110  
Asn Glu Glu Lys Gly Glu Ala Gly Lys Glu Asp Lys Asp Glu Lys Gly  
115 120 125  
Glu Glu Asp Gly Lys Glu Asp Lys Asn Gly Asn Glu Lys Gly Glu Asp  
130 135 140  
Ala Lys Glu Lys Glu Asp Gly Lys Lys Gly Glu Asp Gly Lys Gly Asn  
145 150 155 160  
Gly Glu Asp Gly Lys Glu Lys Gly Glu Asp Glu Lys Glu Glu Glu Asp  
165 170 175  
Arg Lys Glu Thr Gly Val Gly Lys Glu Asn Glu Asp Gly Lys Glu Lys  
180 185 190  
Gly Asp Lys Lys Glu Gly Lys Asp Val Lys Val Lys Glu Asp Glu Lys  
195 200 205  
Glu Arg Glu Asp Gly Lys Glu Asp Glu Gly Gly Asn Glu Glu Glu Ala  
210 215 220  
Gly Lys Glu Lys Glu Asp Leu Lys Glu Glu Glu Gly Lys Glu Glu  
225 230 235 240  
Asp Glu Ile Lys Glu Asp Asp Gly Lys Lys Glu Glu Pro Gln Ser Ile



Lys Glu Asp Leu Lys Glu Glu Glu Gly Lys Glu Glu Asp Glu Ile  
 210 215 220  
 Lys Glu Asp Asp Gly Lys Lys Glu Glu Pro Gln Ser Ile Val  
 225 230 235

<210> 191  
 <211> 699  
 <212> DNA  
 <213> Homo sapiens

<400> 191  
 atgatggaag aaaacataga tacaagtgcc caagcagttg ctgaaaccaa gcaagaagca 60  
 gttgttgaag aagactacaa tgaaaatgct aaaaatggag aagccaaaat tacagaggca 120  
 ccagcttctg aaaaagaaat tgtggaagta aaagaagaaa atattgaaga tgccacagaa 180  
 aagggaggag aaaagaaaga agcagtggca gcagaagtaa aaaatgaaga agaagatcag 240  
 aaagaagatg aagaagatca aaacgaagag aaaggggaag ctggaaaaga agacaaagat 300  
 gaaaaagggg aagaagatgg aaaagaggat aaaaatggaa atgagaaagg agaagatgca 360  
 aaagagaaaag aagatggaaa aaaaggtgaa gacggaaaag gaaatggaga agatggaaaa 420  
 gagaaaggag aagatgaaaa agaggaagaa gacagaaaag aaacaggagt tggaaaagag 480  
 aatgaagatg gaaaagagaa gggagataaa aaagagggga aagatgtaaa agtcaaagaa 540  
 gatgaaaaag agagagaaga tggaaaagaa gatgaaggtg gaaatgagga agaagctgga 600  
 aaagagaaaag aagattttaa agaagaggaa gaagggaaaag aggaagatga gatcaaagaa 660  
 gatgatggaa aaaaagagga gccacagagt attgtttaa 699

<210> 192  
 <211> 232  
 <212> PRT  
 <213> Homo sapiens

<400> 192  
 Met Met Glu Glu Asn Ile Asp Thr Ser Ala Gln Ala Val Ala Glu Thr  
 1 5 10 15  
 Lys Gln Glu Ala Val Val Glu Glu Asp Tyr Asn Glu Asn Ala Lys Asn  
 20 25 30  
 Gly Glu Ala Lys Ile Thr Glu Ala Pro Ala Ser Glu Lys Glu Ile Val  
 35 40 45  
 Glu Val Lys Glu Glu Asn Ile Glu Asp Ala Thr Glu Lys Gly Gly Glu  
 50 55 60  
 Lys Lys Glu Ala Val Ala Ala Glu Val Lys Asn Glu Glu Glu Asp Gln  
 65 70 75 80  
 Lys Glu Asp Glu Glu Asp Gln Asn Glu Glu Lys Gly Glu Ala Gly Lys  
 85 90 95  
 Glu Asp Lys Asp Glu Lys Gly Glu Glu Asp Gly Lys Glu Asp Lys Asn  
 100 105 110  
 Gly Asn Glu Lys Gly Glu Asp Ala Lys Glu Lys Glu Asp Gly Lys Lys  
 115 120 125  
 Gly Glu Asp Gly Lys Gly Asn Gly Glu Asp Gly Lys Glu Lys Gly Glu  
 130 135 140  
 Asp Glu Lys Glu Glu Glu Asp Arg Lys Glu Thr Gly Val Gly Lys Glu  
 145 150 155 160  
 Asn Glu Asp Gly Lys Glu Lys Gly Asp Lys Lys Glu Gly Lys Asp Val  
 165 170 175  
 Lys Val Lys Glu Asp Glu Lys Glu Arg Glu Asp Gly Lys Glu Asp Glu  
 180 185 190  
 Gly Gly Asn Glu Glu Glu Ala Gly Lys Glu Lys Glu Asp Leu Lys Glu

195 200 205  
 Glu Glu Glu Gly Lys Glu Glu Asp Glu Ile Lys Glu Asp Asp Gly Lys  
 210 215 220  
 Lys Glu Glu Pro Gln Ser Ile Val  
 225 230

<210> 193  
 <211> 696  
 <212> DNA  
 <213> Homo sapiens

<400> 193  
 atggaagaaa acatagatac aagtgcccaa gcagttgctg aaaccaagca agaagcagtt 60  
 gttgaagaag actacaatga aaatgctaaa aatggagaag ccaaaattac agaggcacca 120  
 gcttctgaaa aagaaattgt ggaagtaaaa gaagaaaata ttgaagatgc cacagaaaag 180  
 ggaggagaaa agaaagaagc agtggcagca gaagtaaaaa atgaagaaga agatcagaaa 240  
 gaagatgaag aagatcaaaa cgaagagaaa ggggaagctg gaaaagaaga caaagatgaa 300  
 aaaggggaag aagatggaaa agaggataaaa aatggaaatg agaaaggaga agatgcaaaa 360  
 gagaaagaag atggaaaaaa aggtgaagac ggaaaaggaa atggagaaga tggaaaagag 420  
 aaaggagaag atgaaaaaga ggaagaagac agaaaagaaa caggagttgg aaaagagaat 480  
 gaagatggaa aagagaaggg agataaaaaa gaggggaaaag atgtaaaagt caaagaagat 540  
 gaaaaagaga gagaagatgg aaaagaagat gaaggtggaa atgaggaaga agctggaaaa 600  
 gagaaagaag atttaaaaaga agaggaagaa ggaaaagagg aagatgagat caaagaagat 660  
 gatggaaaaa aagaggagcc acagagtatt gtttaa 696

<210> 194  
 <211> 231  
 <212> PRT  
 <213> Homo sapiens

<400> 194  
 Met Glu Glu Asn Ile Asp Thr Ser Ala Gln Ala Val Ala Glu Thr Lys  
 1 5 10 15  
 Gln Glu Ala Val Val Glu Glu Asp Tyr Asn Glu Asn Ala Lys Asn Gly  
 20 25 30  
 Glu Ala Lys Ile Thr Glu Ala Pro Ala Ser Glu Lys Glu Ile Val Glu  
 35 40 45  
 Val Lys Glu Glu Asn Ile Glu Asp Ala Thr Glu Lys Gly Gly Glu Lys  
 50 55 60  
 Lys Glu Ala Val Ala Ala Glu Val Lys Asn Glu Glu Glu Asp Gln Lys  
 65 70 75 80  
 Glu Asp Glu Glu Asp Gln Asn Glu Glu Lys Gly Glu Ala Gly Lys Glu  
 85 90 95  
 Asp Lys Asp Glu Lys Gly Glu Glu Asp Gly Lys Glu Asp Lys Asn Gly  
 100 105 110  
 Asn Glu Lys Gly Glu Asp Ala Lys Glu Lys Glu Asp Gly Lys Lys Gly  
 115 120 125  
 Glu Asp Gly Lys Gly Asn Gly Glu Asp Gly Lys Glu Lys Gly Glu Asp  
 130 135 140  
 Glu Lys Glu Glu Glu Asp Arg Lys Glu Thr Gly Val Gly Lys Glu Asn  
 145 150 155 160  
 Glu Asp Gly Lys Glu Lys Gly Asp Lys Lys Glu Gly Lys Asp Val Lys  
 165 170 175  
 Val Lys Glu Asp Glu Lys Glu Arg Glu Asp Gly Lys Glu Asp Glu Gly  
 180 185 190



<210> 200  
 <211> 18  
 <212> PRT  
 <213> Homo sapiens

<400> 200  
 Met Glu Lys Pro Lys Leu Gln Arg His Gln Leu Leu Lys Lys Lys Leu  
 1 5 10 15  
 Trp Lys

<210> 201  
 <211> 51  
 <212> DNA  
 <213> Homo sapiens

<400> 201  
 atgccacaga aaagggagga gaaaagaaag aagcagtggc agcagaagta a 51

<210> 202  
 <211> 16  
 <212> PRT  
 <213> Homo sapiens

<400> 202  
 Met Pro Gln Lys Arg Glu Glu Lys Arg Lys Lys Gln Trp Gln Gln Lys  
 1 5 10 15

<210> 203  
 <211> 306  
 <212> DNA  
 <213> Homo sapiens

<400> 203  
 atgaagaaga agatcagaaa gaagatgaag aagatcaaaa cgaagagaaa ggggaagctg 60  
 gaaaagaaga caaagatgaa aaaggggaag aagatggaaa agaggataaa aatggaaatg 120  
 agaaaggaga agatgcaaaa gagaaagaag atggaaaaaa aggtgaagac ggaaaaggaa 180  
 atggagaaga tggaaaagag aaaggagaag atgaaaaaga ggaagaagac agaaaagaaa 240  
 caggagtggg aaaagagaat gaagatggaa aagagaaggg agataaaaaa gaggggaaag 300  
 atgtaa 306

<210> 204  
 <211> 101  
 <212> PRT  
 <213> Homo sapiens

<400> 204  
 Met Lys Lys Lys Ile Arg Lys Lys Met Lys Lys Ile Lys Thr Lys Arg  
 1 5 10 15  
 Lys Gly Lys Leu Glu Lys Lys Thr Lys Met Lys Lys Gly Lys Lys Met  
 20 25 30  
 Glu Lys Arg Ile Lys Met Glu Met Arg Lys Glu Lys Met Gln Lys Arg  
 35 40 45  
 Lys Lys Met Glu Lys Lys Val Lys Thr Glu Lys Glu Met Glu Lys Met  
 50 55 60

Glu Lys Arg Lys Glu Lys Met Lys Lys Arg Lys Lys Thr Glu Lys Lys  
 65 70 75 80  
 Gln Glu Leu Glu Lys Arg Met Lys Met Glu Lys Arg Arg Glu Ile Lys  
 85 90 95  
 Lys Arg Gly Lys Met  
 100

<210> 205  
 <211> 282  
 <212> DNA  
 <213> Homo sapiens

<400> 205  
 atgaagaaga tcaaaacgaa gagaaagggg aagctggaaa agaagacaaa gatgaaaaag 60  
 gggaagaaga tggaaaagag gataaaaatg gaaatgagaa aggagaagat gcaaaagaga 120  
 aagaagatgg aaaaaaaggt gaagacggaa aaggaaatgg agaagatgga aaagagaaaag 180  
 gagaagatga aaaagaggaa gaagacagaa aagaaacagg agttggaaaa gagaatgaag 240  
 atggaaaaga gaaggagat aaaaaagagg ggaaagatgt aa 282

<210> 206  
 <211> 93  
 <212> PRT  
 <213> Homo sapiens

<400> 206  
 Met Lys Lys Ile Lys Thr Lys Arg Lys Gly Lys Lys Leu Glu Lys Lys Thr  
 1 5 10 15  
 Lys Met Lys Lys Gly Lys Lys Met Glu Lys Arg Ile Lys Met Glu Met  
 20 25 30  
 Arg Lys Glu Lys Met Gln Lys Arg Lys Lys Met Glu Lys Lys Val Lys  
 35 40 45  
 Thr Glu Lys Glu Met Glu Lys Met Glu Lys Arg Lys Glu Lys Met Lys  
 50 55 60  
 Lys Arg Lys Lys Thr Glu Lys Lys Gln Glu Leu Glu Lys Arg Met Lys  
 65 70 75 80  
 Met Glu Lys Arg Arg Glu Ile Lys Lys Arg Gly Lys Met  
 85 90

<210> 207  
 <211> 231  
 <212> DNA  
 <213> Homo sapiens

<400> 207  
 atgaaaaagg ggaagaagat ggaaaagagg ataaaaatgg aaatgagaaa ggagaagatg 60  
 caaaagagaa agaagatgga aaaaaaggtg aagacggaaa aggaaatgga gaagatggaa 120  
 aagagaaaagg agaagatgaa aaagaggaag aagacagaaa agaaacagga gttggaaaag 180  
 agaatgaaga tggaaaagag aaggagata aaaaagaggg gaaagatgta a 231

<210> 208  
 <211> 76  
 <212> PRT  
 <213> Homo sapiens

<400> 208





<400> 212

Met Glu Met Arg Lys Glu Lys Met Gln Lys Arg Lys Lys Met Glu Lys  
1 5 10 15  
Lys Val Lys Thr Glu Lys Glu Met Glu Lys Met Glu Lys Arg Lys Glu  
20 25 30  
Lys Met Lys Lys Arg Lys Lys Thr Glu Lys Lys Gln Glu Leu Glu Lys  
35 40 45  
Arg Met Lys Met Glu Lys Arg Arg Glu Ile Lys Lys Arg Gly Lys Met  
50 55 60

$\langle 210 \rangle$  213

<211> 189

<212> DNA

<213> Homo sapiens

<400> 213

atgagaaaagg	agaagatgca	aaagagaaaag	aagatggaaa	aaaaggtgaa	gacggaaaag	60
gaaatggaga	agatggaaaa	gagaaaaggag	aagatgaaaa	agaggaagaa	gacagaaaag	120
aaacaggagt	tggaaaagag	aatgaagatg	gaaaagagaa	gggagataaa	aaagagggga	180
aagatgtaa						189

<210> 214

<211> 62

<212> PRT

<213> Homo sapiens

<400> 214

Met	Arg	Lys	Glu	Lys	Met	Gln	Lys	Arg	Lys	Lys	Met	Glu	Lys	Lys	Val
1					5				10					15	
Lys	Thr	Glu	Lys	Glu	Met	Glu	Lys	Met	Glu	Lys	Arg	Lys	Glu	Lys	Met
			20					25					30		
Lys	Lys	Arg	Lys	Lys	Thr	Glu	Lys	Lys	Gln	Glu	Leu	Glu	Lys	Arg	Met
		35					40					45			
Lys	Met	Glu	Lys	Arg	Arg	Glu	Ile	Lys	Lys	Arg	Gly	Lys	Met		
	50					55					60				

<210> 215

<211> 174

<212> DNA

<213> Homo sapiens

<400> 215

atgcaaaaaga	gaaagaagat	ggaaaaaaaag	gtgaagacgg	aaaaggaaat	ggagaagatg	60
tgaaagaagaa	aggagaagat	gaaaaagagg	aagaagacag	aaaagaaaca	ggagttggaa	120
aagagaatga	agatggaaaa	gagaagggag	ataaaaaaga	ggggaaagat	gtaa	174

<210> 216

<211> 57

<212> PRT

<213> Homo sapiens

<400> 216

Met Gln Lys Arg Lys Lys Met Glu Lys Lys Val Lys Thr Glu Lys Glu  
1 5 10 15  
Met Glu Lys Met Glu Lys Arg Lys Glu Lys Met Lys Lys Arg Lys Lys



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 aagagaaaaga agatttaa 78

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Phe Lys Thr Ala Leu Cys Ser Phe Ile Ile Trp  
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<210> 239  
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 atttggttaa 69

<210> 240  
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<400> 240  
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 1 5 10 15  
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Met Tyr Leu His Val Val Lys Leu Ile Glu Ile Asn Ile Phe Ile Lys  
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<212> DNA

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ctttcaactt tgtga 75

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<211> 24

<212> PRT

<213> Homo sapiens

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Met Lys His Leu Ser Ile Asn Phe Val Ile Ile Val Val Glu Tyr Ile  
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<212> DNA

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<211> 5

<212> PRT

<213> Homo sapiens

<400> 248

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<400> 249

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<210> 250

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aaattgtatt cccattttta a 81

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<400> 252  
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Met Lys Leu Tyr Ser His Phe  
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<210> 255  
<211> 27  
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<210> 256  
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ccgagattca aatctccgat ttcccatattg ggggcaagtt tttttcttca ccttcaatat      540
gagaattcag cgaacttgaa agaaaaatca tctgtgagtt ccttcagggtt ctcaactcata      600
gtcatgatcc ttcagaggga atatgcactg gcgagtttaa agtaagggtt atgatatttg      660
atggtcccaa agtacggcag ctgcaaaaag tagtggaagg aaattgtcta cgtgtcttgg      720
aaaaattagt taggaatttg gatgggtaaa aggtaccctt gccttactcc atcttatattt      780
cttagccccc tttgagtgtt ttaactgggtt tcatgtccta gtaggaagtg cattctccat      840
cctcatcctc tgccctccca ggaagtcagt gattgtcttt ttgggcttcc cctccaaagg      900
accttctgca gtggaagtgc cacatccagt tcttttcttt tgttgctgct gtgtttagat      960
aattgaagag atctttgtgc cacacaggat tttttttttt ttttaagaaa aacctataga     1020
tgaaaaatta ctaatgaaac tgtgtgtacg tgtctgtgcg tgcaacataa aaatacagta     1080
gcacctaagg agcttgaatc ttggttcctg taaaatttca aattgatgtg gtattaataa     1140
aaaaaaaaaa aacccaaaaa aaaaaaaaaa aaaagggcgg cc                          1182

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 <213> Homo sapiens

<400> 262  
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 Met Pro Ser Asp Lys Lys Glu  
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<210> 264  
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 <212> DNA  
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 ccactatccc caggaagga aaggtccgc catttgaggaa agtggtttct acgtcactgg 120  
 acaccggttc tgagcattag tttgagaact cgttcccgaa tgtgctttcc tccctctccc 180  
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 35 40 45

Arg Thr Arg Ser Arg Met Cys Phe Pro Pro Ser Pro Leu Pro Thr Ser  
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 <211> 64  
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 Arg Leu Arg His Leu Gly Lys Trp Phe Leu Arg His Trp Thr Pro Val  
 20 25 30  
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 35 40 45  
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 ttactataa 69

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 Val Val Leu Phe Leu  
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 <212> DNA  
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<211> 6  
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Met Arg Ile Gln Arg Thr  
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<210> 276  
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54

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<210> 287
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aggcttccta aggggaagact tcctgtccca aaggaagtga accgcaagaa gaacgatgag 240  
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120

132

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			20				25						30		
Gln	Trp	Thr	Asn	Arg	Cys	His	Gln	Thr	Ser	Phe					
			35				40								

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<211> 33

<212> DNA

<213> Homo sapiens

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<212> PRT

<213> Homo sapiens

<400> 306

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<211> 36

<212> DNA

<213> Homo sapiens











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Glu	Ser	Leu	Phe	Ala	Cys	Asn	Ile	Cys	Pro	Tyr	Val	Val	Leu	Asp	Gly
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Arg	Glu	Lys	Ile	Gln	Met	Ala	His	Ser	Leu	Ser	Val	Gly	Gly	Ser	Gly
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Tyr	Val	Cys	Pro	Leu	Leu	Ile	Arg	Glu	Val	Phe	Ile	Gln	Val	Leu	Ile
			85						90					95	
Lys	Leu	Arg	Val	Cys	Phe	Val	Gln	Cys	Phe	Ser	Glu	Ala	Asp	Arg	Asp
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			165						170					175	
Asn	Met	Asn	Lys	Ala	Leu	Leu	Pro	Leu	Phe	Ala	Val	Leu	Cys	Gly	Asn
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		210				215						220			
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225				230						235					240
Leu	Asp	Asn	Val	Leu	Lys	Tyr	Leu	Pro	Lys	Lys	Asp	Arg	Glu	Asn	Val
			245							250				255	
Lys	Glu	Leu	Leu	Cys	Cys	Ser	Met	Glu	Glu	Tyr	Gln	Gln	Ser	Gln	Val
		260						265					270		
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Phe	Val	Gln	Cys	Phe	Ser	Glu	Ala	Asp	Arg	Asp	Ile	Met	Thr	Leu	Ala
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Phe	Asp	Leu	Lys	Thr	Gly	Phe	Cys	Pro	Leu	Asn	Ser	Phe	Gln	Trp	Arg
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Asn	Met	Asn	Thr	Ile	Lys	Gly	Thr	Gln	Asn	Tyr	Ile	Pro	Ala	Lys	Cys
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Trp	Leu	Ser	His	Phe	Ala	Asn	Pro	Thr	Glu	Ala	Leu	Asp	Asn	Val	Leu
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Lys	Tyr	Leu	Pro	Lys	Lys	Asp	Arg	Glu	Asn	Val	Lys	Glu	Leu	Leu	Cys
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Cys	Ser	Met	Glu	Glu	Tyr	Gln	Gln	Ser	Gln	Val	Lys	Leu	Gln	Asp	Phe
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Gln	Ile	Ile	Tyr	Gly	Leu	Leu	Leu	Asn	Ala	Ser	Pro	His	Leu	Asp	Lys
		275					280					285			
Thr	Ser	Trp	Asn	Ala	Leu	Pro	Pro	Gln	Pro	Leu	Ala	Phe	Ser	Glu	Val
	290					295					300				
Glu	Arg	Ile	Asn	Lys	Asn	Ile	Arg	Thr	Ser	Ile	Ile	Asp	Ala	Val	Glu
305					310					315					320
Leu	Ala	Lys	Asp	His	Ser	Asp	Leu	Ser	Arg	Leu	Thr	Glu	Leu	Ser	Leu
				325					330					335	
Arg	Arg	Arg	Gln	Met	Leu	Leu	Leu	Glu	Thr	Leu	Lys	Val	Lys	Gln	Thr
			340					345					350		
Ile	Leu	Glu	Pro	Ile	Pro	Thr	Ser	Leu	Lys	Leu	Pro	Ile	Ala	Val	Ser
		355					360					365			
Cys	Tyr	Trp	Leu	Gln	His	Thr	Glu	Thr	Lys	Ala	Lys	Leu	His	His	Leu
	370					375					380				
Gln	Ser	Leu	Leu	Leu	Thr	Met	Leu	Val	Gly	Pro	Leu	Ile	Ala	Ile	Ile
385					390					395					400

Asn Ser Pro Gly Asn Val Asp Pro Val Pro Arg Gln Ala Gln Cys Leu  
 405 410 415  
 Ala Pro Arg

<210> 390  
 <211> 45  
 <212> DNA  
 <213> Homo sapiens

<400> 390  
 atgtatgtcc cttactcatc cgggaagtat tcatacaggt ttgga 45

<210> 391  
 <211> 14  
 <212> PRT  
 <213> Homo sapiens

<400> 391  
 Met Tyr Val Pro Tyr Ser Ser Gly Lys Tyr Ser Tyr Arg Phe  
 1 5 10

<210> 392  
 <211> 108  
 <212> DNA  
 <213> Homo sapiens

<400> 392  
 atgtccctta ctcacccggg aagtattcat acagggttttg atcaagctgc ggggtgtgttt 60  
 tgtccagtgc ttttcagaag cagatcggga cattatgaca cttgctaa 108

<210> 393  
 <211> 35  
 <212> PRT  
 <213> Homo sapiens

<400> 393  
 Met Ser Leu Thr His Pro Gly Ser Ile His Thr Gly Phe Asp Gln Ala  
 1 5 10 15  
 Ala Gly Val Phe Cys Pro Val Leu Phe Arg Ser Arg Ser Gly His Tyr  
 20 25 30  
 Asp Thr Cys  
 35

<210> 394  
 <211> 1128  
 <212> DNA  
 <213> Homo sapiens

<400> 394  
 atgacacttg ctaaccattg gaattgccct gtgttatcat cagatagtga cttttgcatt 60  
 tttgacctga aaactggggt ttgccattg aatagctttc agtggagaaa tatgaacact 120  
 attaagggca cacaaaacta tatccctgcc aaatgctttt cccttgatgc attctgccat 180  
 cacttcagca atatgaataa agctctacta cctctctttg cggtgctatg tggaaatgac 240  
 catgttaatc taccatcat ggagacattc ttaagtaaag cgcgtcttcc tcttggagct 300





Asp Ala Val Glu Leu Ala Lys Asp His Ser Asp Leu Ser Arg Leu Thr  
           275                                  280                                  285  
 Glu Leu Ser Leu Arg Arg Arg Gln Met Leu Leu Leu Glu Thr Leu Lys  
           290                                  295                                  300  
 Val Lys Gln Thr Ile Leu Glu Pro Ile Pro Thr Ser Leu Lys Leu Pro  
 305                                  310                                  315                                  320  
 Ile Ala Val Ser Cys Tyr Trp Leu Gln His Thr Glu Thr Lys Ala Lys  
                                   325                                  330                                  335  
 Leu His His Leu Gln Ser Leu Leu Leu Thr Met Leu Val Gly Pro Leu  
                                   340                                  345                                  350  
 Ile Ala Ile Ile Asn Ser Pro Gly Asn Val Asp Pro Val Pro Arg Gln  
                                   355                                  360                                  365  
 Ala Gln Cys Leu Ala Pro Arg  
                                   370                                  375

<210> 396

<211> 1017

<212> DNA

<213> Homo sapiens

<400> 396

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ggaaatgacc	atgttaatct	acccatcatg	gagacattct	taagtaaagc	gcgtcttcct	180
cttgaggcta	ccagttctaa	agggaggaga	caccaccgaa	tcctgggact	tctgaattgg	240
ttgtctcatt	ttgccaaccc	taccgaagca	ctagataatg	ttctgaaata	cctcccaaaa	300
aaggatcgag	aaaatgttaa	ggaacttctc	tgctgttcca	tggaagaata	ccaacagtcc	360
caggtgaagc	tacaggactt	cttcagtggt	ggtacttatg	tctgtccaga	tgcttgaat	420
cttggtttac	cagaatgggt	attagtggct	ttagctaaag	gccagctatc	tcctttcatc	480
agtgatgctt	tggtcctaag	acggaccatt	cttccacac	aggtggaaaa	catgcagcaa	540
ccaaatgccc	acagaatatc	tcagcccatc	aggcaaata	tctatgggct	tccttttaaat	600
gcctcaccac	atctggacaa	gacatcctgg	aatgcattgc	ctcctcagcc	tctagctttc	660
agtgaagtgg	aaaggattaa	taaaaatatc	agaacctcaa	tcattgatgc	agtagaactg	720
gccaaagatc	attctgactt	aagcagattg	actgagctct	ccttgaggag	gcggcagatg	780
cttctgttag	aaacctgaa	ggtgaaacag	accattctgg	agccaatccc	tacttcactg	840
aagttgccca	ttgctgtcag	ttgctactgg	ttgcagcaca	ccgagaccaa	agcaaagcta	900
catcatctac	aatccttact	gctcacaatg	ctagtggggc	ccttgattgc	cataatcaac	960
agccctggaa	atgtggaccc	tgtaccagg	caggctcagt	gtcttgctcc	tcgctag	1017

<210> 397

<211> 338

<212> PRT

<213> Homo sapiens

<400> 397

Met	Asn	Thr	Ile	Lys	Gly	Thr	Gln	Asn	Tyr	Ile	Pro	Ala	Lys	Cys	Phe
1				5					10					15	
Ser	Leu	Asp	Ala	Phe	Cys	His	His	Phe	Ser	Asn	Met	Asn	Lys	Ala	Leu
			20					25					30		
Leu	Pro	Leu	Phe	Ala	Val	Leu	Cys	Gly	Asn	Asp	His	Val	Asn	Leu	Pro
			35				40					45			
Ile	Met	Glu	Thr	Phe	Leu	Ser	Lys	Ala	Arg	Leu	Pro	Leu	Gly	Ala	Thr
	50					55				60					
Ser	Ser	Lys	Gly	Arg	Arg	His	His	Arg	Ile	Leu	Gly	Leu	Leu	Asn	Trp
65					70				75					80	







<211> 12  
 <212> PRT  
 <213> Homo sapiens

<400> 407  
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 1 5 10

<210> 408  
 <211> 33  
 <212> DNA  
 <213> Homo sapiens

<400> 408  
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33

<210> 409  
 <211> 10  
 <212> PRT  
 <213> Homo sapiens

<400> 409  
 Met Leu Ile Tyr Pro Ser Trp Arg His Ser  
 1 5 10

<210> 410  
 <211> 870  
 <212> DNA  
 <213> Homo sapiens

<400> 410  
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 gcactagata atgttctgaa atacctccca aaaaaggatc gagaaaatgt taaggaactt 180  
 ctctgctggt ccatggaaga ataccaacag tcccagggtga agctacagga cttcttccag 240  
 tgtggtactt atgtctgtcc agatgccttg aatcttgggt taccagaatg ggtattagt 300  
 gcttttagcta aaggccagct atctcctttc atcagtgatg ctttggtcct aagacggacc 360  
 attcttccca cacagggtgga aaacatgcag caaccaaagt cccacagaat atctcagccc 420  
 atcaggcaaa tcactctatgg gcttctttta aatgcctcac cacatctgga caagacatcc 480  
 tggaaatgcat tgcctcctca gcctctagct ttcagtgaag tggaaaggat taataaaaaat 540  
 atcagaacct caatcattga tgcagtagaa ctggccaagg atcattctga ctttaagcaga 600  
 ttgactgagc tctccttgag gaggcggcag atgcttctgt tagaaaccct gaaggtgaaa 660  
 cagaccattc tggagccaat ccctacttca ctgaagttgc ccattgctgt cagttgctac 720  
 tggttgcagc acaccgagac caaagcaaag ctacatcatc tacaatcctt actgctcaca 780  
 atgctagtgg ggcccttgat tgccataatc aacagccctg gaaatgtgga ccctgtaccc 840  
 aggcaggctc agtgtcttgc tcctcgctag 870

<210> 411  
 <211> 289  
 <212> PRT  
 <213> Homo sapiens

<400> 411  
 Met Glu Thr Phe Leu Ser Lys Ala Arg Leu Pro Leu Gly Ala Thr Ser  
 1 5 10 15

Ser Lys Gly Arg Arg His His Arg Ile Leu Gly Leu Leu Asn Trp Leu  
20 25 30  
Ser His Phe Ala Asn Pro Thr Glu Ala Leu Asp Asn Val Leu Lys Tyr  
35 40 45  
Leu Pro Lys Lys Asp Arg Glu Asn Val Lys Glu Leu Leu Cys Cys Ser  
50 55 60  
Met Glu Glu Tyr Gln Gln Ser Gln Val Lys Leu Gln Asp Phe Phe Gln  
65 70 75 80  
Cys Gly Thr Tyr Val Cys Pro Asp Ala Leu Asn Leu Gly Leu Pro Glu  
85 90 95  
Trp Val Leu Val Ala Leu Ala Lys Gly Gln Leu Ser Pro Phe Ile Ser  
100 105 110  
Asp Ala Leu Val Leu Arg Arg Thr Ile Leu Pro Thr Gln Val Glu Asn  
115 120 125  
Met Gln Gln Pro Asn Ala His Arg Ile Ser Gln Pro Ile Arg Gln Ile  
130 135 140  
Ile Tyr Gly Leu Leu Leu Asn Ala Ser Pro His Leu Asp Lys Thr Ser  
145 150 155 160  
Trp Asn Ala Leu Pro Pro Gln Pro Leu Ala Phe Ser Glu Val Glu Arg  
165 170 175  
Ile Asn Lys Asn Ile Arg Thr Ser Ile Ile Asp Ala Val Glu Leu Ala  
180 185 190  
Lys Asp His Ser Asp Leu Ser Arg Leu Thr Glu Leu Ser Leu Arg Arg  
195 200 205  
Arg Gln Met Leu Leu Leu Glu Thr Leu Lys Val Lys Gln Thr Ile Leu  
210 215 220  
Glu Pro Ile Pro Thr Ser Leu Lys Leu Pro Ile Ala Val Ser Cys Tyr  
225 230 235 240  
Trp Leu Gln His Thr Glu Thr Lys Ala Lys Leu His His Leu Gln Ser  
245 250 255  
Leu Leu Leu Thr Met Leu Val Gly Pro Leu Ile Ala Ile Ile Asn Ser  
260 265 270  
Pro Gly Asn Val Asp Pro Val Pro Arg Gln Ala Gln Cys Leu Ala Pro  
275 280 285  
Arg

<210> 412  
<211> 54  
<212> DNA  
<213> Homo sapiens

<400> 412  
atgttaagga acttctctgc tgttccatgg aagaatacca acagtcccag gtga

54

<210> 413  
<211> 17  
<212> PRT  
<213> Homo sapiens

<400> 413  
Met Leu Arg Asn Phe Ser Ala Val Pro Trp Lys Asn Thr Asn Ser Pro  
1 5 10 15  
Arg

<210> 414  
 <211> 678  
 <212> DNA  
 <213> Homo sapiens

<400> 414  
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 gtctgtccag atgccttgaa tcttggttta ccagaatggg tattagtggc tttagctaaa 120  
 ggccagctat ctcctttcat cagtgatgct ttggtcctaa gacggaccat tcttcccaca 180  
 cagggtgaaa acatgcagca accaaatgcc cacagaatat ctcagcccat caggcaaatc 240  
 atctatgggc ttctttttaa tgccctacca catctggaca agacatcctg gaatgcattg 300  
 cctcctcagc ctctagcttt cagtgaagtg gaaaggatta ataaaaatat cagaacctca 360  
 atcattgatg cagtagaact ggccaaggat cattctgact taagcagatt gactgagctc 420  
 tccttgagga ggcggcagat gcttctgtta gaaaccctga aggtgaaaca gaccattctg 480  
 gagccaatcc ctacttcact gaagttgccc attgctgtca gttgctactg gttgcagcac 540  
 accgagacca aagcaaagct acatcatcta caatccttac tgctcacaat gctagtgggg 600  
 cccttgattg ccataatcaa cagccctgga aatgtggacc ctgtaccacag gcaggctcag 660  
 tgtcttgctc ctcgctag 678

<210> 415  
 <211> 225  
 <212> PRT  
 <213> Homo sapiens

<400> 415  
 Met Glu Glu Tyr Gln Gln Ser Gln Val Lys Leu Gln Asp Phe Phe Gln  
 1 5 10 15  
 Cys Gly Thr Tyr Val Cys Pro Asp Ala Leu Asn Leu Gly Leu Pro Glu  
 20 25 30  
 Trp Val Leu Val Ala Leu Ala Lys Gly Gln Leu Ser Pro Phe Ile Ser  
 35 40 45  
 Asp Ala Leu Val Leu Arg Arg Thr Ile Leu Pro Thr Gln Val Glu Asn  
 50 55 60  
 Met Gln Gln Pro Asn Ala His Arg Ile Ser Gln Pro Ile Arg Gln Ile  
 65 70 75 80  
 Ile Tyr Gly Leu Leu Asn Ala Ser Pro His Leu Asp Lys Thr Ser  
 85 90 95  
 Trp Asn Ala Leu Pro Pro Gln Pro Leu Ala Phe Ser Glu Val Glu Arg  
 100 105 110  
 Ile Asn Lys Asn Ile Arg Thr Ser Ile Ile Asp Ala Val Glu Leu Ala  
 115 120 125  
 Lys Asp His Ser Asp Leu Ser Arg Leu Thr Glu Leu Ser Leu Arg Arg  
 130 135 140  
 Arg Gln Met Leu Leu Leu Glu Thr Leu Lys Val Lys Gln Thr Ile Leu  
 145 150 155 160  
 Glu Pro Ile Pro Thr Ser Leu Lys Leu Pro Ile Ala Val Ser Cys Tyr  
 165 170 175  
 Trp Leu Gln His Thr Glu Thr Lys Ala Lys Leu His His Leu Gln Ser  
 180 185 190  
 Leu Leu Leu Thr Met Leu Val Gly Pro Leu Ile Ala Ile Ile Asn Ser  
 195 200 205  
 Pro Gly Asn Val Asp Pro Val Pro Arg Gln Ala Gln Cys Leu Ala Pro  
 210 215 220  
 Arg  
 225





<400> 422  
atgcagcaac caaatgccca cagaatatct cagcccatca ggcaaatacat ctatgggctt 60  
cttttaaatg cctcaccaca tctggacaag acatcctgga atgcattgcc tcttcagcct 120  
ctagctttca gtgaagtgga aaggattaat aaaaatatca gaacctcaat cattgatgca 180  
gtagaactgg ccaaggatca ttctgactta agcagattga ctgagctctc cttgaggagg 240  
cggcagatgc ttctgttaga aacctgaag gtgaaacaga ccattctgga gccaatccct 300  
acttactga agttgcccat tgctgtcagt tgctactggt tgcagcacac cgagacaaaa 360  
gcaaagctac atcatctaca atccttactg ctcacaatgc tagtggggcc cttgattgcc 420  
ataatcaaca gccctggaaa tgtggaccct gtaccaggc aggctcagtg tcttgctcct 480  
cgctag 486

<210> 423  
<211> 161  
<212> PRT  
<213> Homo sapiens

<400> 423  
Met Gln Gln Pro Asn Ala His Arg Ile Ser Gln Pro Ile Arg Gln Ile  
1 5 10 15  
Ile Tyr Gly Leu Leu Leu Asn Ala Ser Pro His Leu Asp Lys Thr Ser  
20 25 30  
Trp Asn Ala Leu Pro Pro Gln Pro Leu Ala Phe Ser Glu Val Glu Arg  
35 40 45  
Ile Asn Lys Asn Ile Arg Thr Ser Ile Ile Asp Ala Val Glu Leu Ala  
50 55 60  
Lys Asp His Ser Asp Leu Ser Arg Leu Thr Glu Leu Ser Leu Arg Arg  
65 70 75 80  
Arg Gln Met Leu Leu Leu Glu Thr Leu Lys Val Lys Gln Thr Ile Leu  
85 90 95  
Glu Pro Ile Pro Thr Ser Leu Lys Leu Pro Ile Ala Val Ser Cys Tyr  
100 105 110  
Trp Leu Gln His Thr Glu Thr Lys Ala Lys Leu His His Leu Gln Ser  
115 120 125  
Leu Leu Leu Thr Met Leu Val Gly Pro Leu Ile Ala Ile Ile Asn Ser  
130 135 140  
Pro Gly Asn Val Asp Pro Val Pro Arg Gln Ala Gln Cys Leu Ala Pro  
145 150 155 160  
Arg

<210> 424  
<211> 54  
<212> DNA  
<213> Homo sapiens

<400> 424  
atgccacag aatatctcag cccatcaggc aaatcatcta tgggcttctt ttaa 54

<210> 425  
<211> 17  
<212> PRT  
<213> Homo sapiens

<400> 425  
Met Pro Thr Glu Tyr Leu Ser Pro Ser Gly Lys Ser Ser Met Gly Phe

15

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<210> 426
<211> 15
<212> DNA
<213> Homo sapiens
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<400> 426  
atgggcttct tttaa

15

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<210> 427
<211> 4
<212> PRT
<213> Homo sapiens
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<400> 427  
Met Gly Phe Phe  
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<210> 428
<211> 57
<212> DNA
<213> Homo sapiens
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<400> 428  
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57

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<210> 429
<211> 18
<212> PRT
<213> Homo sapiens
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```
<400> 429
Met Pro His His Ile Trp Thr Arg His Pro Gly Met His Cys Leu Leu
 1           5           10           15
Ser Leu
```

```
<210> 430
<211> 24
<212> DNA
<213> Homo sapiens
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<400> 430
atgcattgcc tcctcagcct ctag
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24

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<210> 431
<211> 7
<212> PRT
<213> Homo sapiens
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<400> 431  
Met His Cys Leu Leu Ser Leu

1

5

&lt;210&gt; 432

&lt;211&gt; 240

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 432

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atgcttctgt tagaaaccct gaaggtgaaa cagaccattc tggagccaat ccctacttca      60
ctgaagttgc ccattgctgt cagttgctac tggttgcagc acaccgagac caaagcaaag      120
ctacatcatc tacaatcctt actgctcaca atgctagtgg ggcccttgat tgccataatc      180
aacagccctg gaaatgtgga ccctgtaccc aggcaggctc agtgtcttgc tcctcgctag      240

```

&lt;210&gt; 433

&lt;211&gt; 79

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 433

```

Met Leu Leu Leu Glu Thr Leu Lys Val Lys Gln Thr Ile Leu Glu Pro
 1          5          10          15
Ile Pro Thr Ser Leu Lys Leu Pro Ile Ala Val Ser Cys Tyr Trp Leu
      20          25          30
Gln His Thr Glu Thr Lys Ala Lys Leu His His Leu Gln Ser Leu Leu
      35          40          45
Leu Thr Met Leu Val Gly Pro Leu Ile Ala Ile Ile Asn Ser Pro Gly
      50          55          60
Asn Val Asp Pro Val Pro Arg Gln Ala Gln Cys Leu Ala Pro Arg
65          70          75

```

&lt;210&gt; 434

&lt;211&gt; 90

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens

&lt;400&gt; 434

```

atgctagtgg ggcccttgat tgccataatc aacagccctg gaaatgtgga ccctgtaccc      60
aggcaggctc agtgtcttgc tcctcgctag      90

```

&lt;210&gt; 435

&lt;211&gt; 29

&lt;212&gt; PRT

&lt;213&gt; Homo sapiens

&lt;400&gt; 435

```

Met Leu Val Gly Pro Leu Ile Ala Ile Ile Asn Ser Pro Gly Asn Val
 1          5          10          15
Asp Pro Val Pro Arg Gln Ala Gln Cys Leu Ala Pro Arg
      20          25

```

&lt;210&gt; 436

&lt;211&gt; 54

&lt;212&gt; DNA

&lt;213&gt; Homo sapiens





```
<210> 447
<211> 65
<212> PRT
<213> Homo sapiens
```

```
<210> 448
<211> 5'
<212> DNA
<213> Homo sapiens
```

```
<210> 449
<211> 18
<212> PRT
<213> Homo sapiens
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```
<210> 450
<211> 231
<212> DNA
<213> Homo sapiens
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102

agagggagaa ccactgcaca caccaagtgt tggatatgagg gaaacaaccg gtttggggtg 180  
ttaatgggtg aaaacttaga ggaacatagt gaggcctcca acattgaata a 231

<210> 451  
<211> 76  
<212> PRT  
<213> Homo sapiens

<400> 451  
Met Asn Ile Tyr Ser Met Pro Thr Arg Ser Tyr Ala Pro Ala Glu Ile  
1 5 10 15  
Phe Leu Pro Lys Gly Arg Ser Asn Ser Lys Lys Lys Arg Gln Lys Lys  
20 25 30  
Gln Asn Thr Ser Cys Ser Lys Asn Arg Gly Arg Thr Thr Ala His Thr  
35 40 45  
Lys Cys Trp Tyr Glu Gly Asn Asn Arg Phe Gly Leu Leu Met Val Glu  
50 55 60  
Asn Leu Glu Glu His Ser Glu Ala Ser Asn Ile Glu  
65 70 75

<210> 452  
<211> 216  
<212> DNA  
<213> Homo sapiens

<400> 452  
atgccacaaa ggtcatatgc ccccgctgaa atattcctac caaaaggtag atcaaattca 60  
aaaaaaaaaa ggcagaagaa acagaatacc agctgttcta agaacagagg gagaaccact 120  
gcacacacca agtgttggtg tgaggggaaac aaccggtttg ggttggttaat gggtgaaaac 180  
ttagaggaac atagtgaggc ctccaacatt gaataa 216

<210> 453  
<211> 71  
<212> PRT  
<213> Homo sapiens

<400> 453  
Met Pro Thr Arg Ser Tyr Ala Pro Ala Glu Ile Phe Leu Pro Lys Gly  
1 5 10 15  
Arg Ser Asn Ser Lys Lys Lys Arg Gln Lys Lys Gln Asn Thr Ser Cys  
20 25 30  
Ser Lys Asn Arg Gly Arg Thr Thr Ala His Thr Lys Cys Trp Tyr Glu  
35 40 45  
Gly Asn Asn Arg Phe Gly Leu Leu Met Val Glu Asn Leu Glu Glu His  
50 55 60  
Ser Glu Ala Ser Asn Ile Glu  
65 70

<210> 454  
<211> 153  
<212> DNA  
<213> Homo sapiens

<400> 454  
atgccccgcg tgaaatattc ctacaaaaag gtagatcaaa ttcaaaaaaa aaaaggcaga 60

agaaacagaa taccagctgt tctaagaaca gagggagaac cactgcacac accaagtgtt 120  
 ggtatgaggg aaacaaccgg tttgggttgt taa 153

<210> 455  
 <211> 50  
 <212> PRT  
 <213> Homo sapiens

<400> 455  
 Met Pro Pro Leu Lys Tyr Ser Tyr Gln Lys Val Asp Gln Ile Gln Lys  
 1 5 10 15  
 Lys Lys Gly Arg Arg Asn Arg Ile Pro Ala Val Leu Arg Thr Glu Gly  
 20 25 30  
 Glu Pro Leu His Thr Pro Ser Val Gly Met Arg Glu Thr Thr Gly Leu  
 35 40 45  
 Gly Cys  
 50

<210> 456  
 <211> 30  
 <212> DNA  
 <213> Homo sapiens

<400> 456  
 atgagggaaa caaccggttt gggttgttaa 30

<210> 457  
 <211> 9  
 <212> PRT  
 <213> Homo sapiens

<400> 457  
 Met Arg Glu Thr Thr Gly Leu Gly Cys  
 1 5

<210> 458  
 <211> 48  
 <212> DNA  
 <213> Homo sapiens

<400> 458  
 atggttgaaa acttagagga acatagtgag gcotccaaca ttgaataa 48

<210> 459  
 <211> 15  
 <212> PRT  
 <213> Homo sapiens

<400> 459  
 Met Val Glu Asn Leu Glu Glu His Ser Glu Ala Ser Asn Ile Glu  
 1 5 10 15

<210> 460  
 <211> 15  
 <212> DNA



[illegible]

15

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<400> 461
Met Tyr Leu Ile
1
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<400>	462						
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agaagcggaa	gcggaagaag	aacgcccctc	ccgagcccgt	gccgcccccc	cgtgccgccc		120
ccgccccac	ccacgtccgc	tccccgcagc	ccccgcccc	cgcccccgct	cccgcacgag		180
acgagctgcc	ggactggaac	gaggtgctcc	cgccctggga	tcgggaggag	gacgaggtgt		240
accgcgcagg	gccgtaccac	cctttcccca	actacatccg	gccgcggaca	ctgcagccgc		300
cctcggcctt	gcgcgcgcgc	cactaccacc	acgccttgcc	gccttcgcgc	cactatcccg		360
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ccagcccagg agggcggggtg gtttgtgcga gttcccttgc cacgcggggc cccggcccca      180
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tga                                          243
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